

GenCore version 5.1.4_p5_4578
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OM protein - protein search, using sw model

Run on: May 19, 2003, 16:48:18 ; Search time 32.861 Seconds
(without alignments)
1056.640 Million cell updates/sec

Title: US-09-625-573-4
Perfect score: 1900
Sequence: 1 MLTSRSRFRNTNESEEEV.....DGVTSTNTPSTGEQEVSAGL 360

Scoring table: BLOSUM62
Gapext 0.5

Searched: 362588 seqs, 96450795 residues

Total number of hits satisfying chosen parameters: 362588

Minimum DB seq length: 0
Maximum DB seq length: 2000000000
Post-processing: Minimum Match 100%

Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA:
 1: /cgn2_6/ptodata/2/pubpaas/DS08_NEW_PUB.pep:
 2: /cgn2_6/ptodata/2/pubpaas/PCB_PUB.pep:
 3: /cgn2_6/ptodata/2/pubpaas/DS06_NEW_PUB.pep:
 4: /cgn2_6/ptodata/2/pubpaas/DS06_PUBCOMB.pep:
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 8: /cgn2_6/ptodata/2/pubpaas/DS08_PUBCOMB.pep:
 9: /cgn2_6/ptodata/2/pubpaas/DS09_NEW_PUB.pep:
 10: /cgn2_6/ptodata/2/pubpaas/US09_NEW_PUB.pep:
 11: /cgn2_6/ptodata/2/pubpaas/US10_NEW_PUB.pep:
 12: /cgn2_6/ptodata/2/pubpaas/US10_PUBCOMB.pep:
 13: /cgn2_6/ptodata/2/pubpaas/US60_NEW_PUB.pep:
 14: /cgn2_6/ptodata/2/pubpaas/US60_PUBCOMB.pep:
 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	DB ID	Description
1	100.0	360	10 US-09-131-827A-2	Sequence 2, Appli
2	1899.9	360	10 US-09-131-827A-2	Sequence 20, Appli
3	1873.8	360	10 US-09-131-827A-2	Sequence 7, Appli
4	1873.8	360	10 US-09-131-827A-2	Sequence 7, Appli
5	1873.8	360	10 US-09-131-827A-2	Sequence 7, Appli
6	1568.5	82.6	347 9	Sequence 3, Appli
7	1568.5	82.6	347 10	Sequence 9, Appli
8	1568.5	82.6	344 10	Sequence 9, Appli
9	1568.5	82.6	344 10	Sequence 9, Appli
10	1473	77.5	329 10	Sequence 9, Appli
11	1473	77.5	329 10	Sequence 9, Appli
12	1473	77.5	329 10	Sequence 9, Appli
13	1473	77.5	329 10	Sequence 9, Appli
14	1364	71.8	352 10	Sequence 2, Appli
15	1364	71.8	352 10	Sequence 15, Appli
16	1364	71.8	352 10	Sequence 1, Appli
17	1364	71.8	352 10	Sequence 5, Appli
18	1364	71.8	352 10	Sequence 5, Appli
19	1364	71.8	352 10	Sequence 5, Appli

ORGANISM: Homo sapiens

US-09-131-827A-2

RESULT 1
US-09-131-827A-2
Sequence 2, Application US/09131827A
Patent No. US20020038469A1
GENERAL INFORMATION:
APPLICANT: Dean, Michael J.
O'Brien, Stephen J.
Smith, Michael
APPLICANT: Carrington, Mary
APPLICANT: DELAYED PROGRESSION TO AIDS BY A
TITLE OF INVENTION: MISSENSE ALLELE OF THE CCR2 GENE
FILE REFERENCE: 14-014-0333
CURRENT APPLICATION NUMBER: US/09-131, 827A
CURRENT FILING DATE: 1998-08-10
PRIOR APPLICATION NUMBER: 60/055, 659
PRIOR FILING DATE: 1997-08-14
NUMBER OF SEQ ID NOS: 20
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO: 2
LENGTH: 360
TYPE: PRP
ORGANISM: Homo sapiens
US-09-131-827A-2

Query Match 100.0%; Score 1900; DB 10;
Best Local Similarity 100.0%; Pred. No. 2, 8e-16;
Matches 360; Conservative 0; Mismatches 0;
Indels 0; Gaps 0;

QY 1 MLLVILINCKKLRKCLTDYLNAISDLFLPLTLPLWAHSANANEVNAMCKLFGLY 120
Db 1 MLLVILINCKKLRKCLTDYLNAISDLFLPLTLPLWAHSANANEVNAMCKLFGLY 120
QY 61 HIGYFGGIFIFIITIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAFASTPGIIFTK 180
Db 61 HIGYFGGIFIFIITIDRYLAIVHAVFALKARTVTFGVVTSVITWLVAFASTPGIIFTK 180
QY 121 COKEDEYYVCGPFPRGNNNFHTIMRNLLGVLPLUMIVCYSGILKTLRCKNEKKHR 240
QY 181 COKEDEYYVCGPFPRGNNNFHTIMRNLLGVLPLUMIVCYSGILKTLRCKNEKKHR 240

Sequence 2, Appli

Sequence 20, Appli

Sequence 17, Appli

Sequence 2, Appli

Sequence 22, Appli

Sequence 22, Appli

Sequence 2, Appli

Db 181 CQEDSVYVCGPFPRGWNNTIMRNLLGIVLPLIMIVCYSGILKTLRCRNEKKHR 240
 Qy 241 AYRVFTIMIVYFLWTPNIVILLNTQEFFGLSNCESTSOLDOATQVETLGMTMHCCT 300
 Db 241 AYRVFTIMIVYFLWTPNIVILLNTQEFFGLSNCESTSOLDOATQVETLGMTMHCCT 300
 Qy 301 NPIIYAFGEKFRRLSYFFRKHITKRFCKOCPVYRETDGVTSNTNSTGEOEVSAGL 360
 Db 301 NPIIYAFGEKFRRLSYFFRKHITKRFCKOCPVYRETDGVTSNTNSTGEOEVSAGL 360

RESULT 2
 US-09-131-827A-20
 ; Sequence 20, Application US/09131827A
 ; Patent No. US2002008469A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Dean, Michael
 ; APPLICANT: O'Brien, Stephen J.
 ; APPLICANT: Smith, Michael
 ; APPLICANT: Carrington, Mary
 ; TITLE OF INVENTION: DELAYED PROGRESSION TO AIDS BY A
 ; TITLE OF INVENTION: MISSENSE ALLELE OF THE CCR2 GENE
 ; FILE REFERENCE: 14014 0333
 ; CURRENT APPLICATION NUMBER: US/09/131 , 827A
 ; CURRENT FILING DATE: 1998-08-10
 ; PRIOR APPLICATION NUMBER: 60/055, 659
 ; PRIOR FILING DATE: 1997-08-14
 ; NUMBER OF SEQ ID NOS: 20
 ; SOFTWARE: FastSEQ for Windows Version 4.0
 ; SEQ ID NO: 20
 ; LENGTH: 360
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 ; US-09-131-827A-20

Query Match Score 99.9%; Pred. No. 3.5e-163; Length 360;

Best Local Similarity 99.7%;保守性 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 MLSTSRSRFTNTNESEEYTFFDYGACHKFVDKIQGAQLLPPLYSVLFIFGVGN 60
 Db 1 MLSTSRSRFTNTNESEEYTFFDYGACHKFVDKIQGAQLLPPLYSVLFIFGVGN 60
 Qy 61 MLVVLLINCKKLKCUDTIVLNLAISDLLELTPLWAHSANNEWFGNAMCKLFGLY 120
 Db 61 MLVVLLINCKKLKCUDTIVLNLAISDLLELTPLWAHSANNEWFGNAMCKLFGLY 120
 Qy 121 HIGYEGGIFTILLTIDRYLAIVHAWALKARTVTGGVTSVITWLVAFAVGIFTK 180
 Db 121 HIGYEGGIFTILLTIDRYLAIVHAWALKARTVTGGVTSVITWLVAFAVGIFTK 180
 Qy 181 CQEDSVYVCGPFPRGWNNTIMRNLLGIVLPLIMIVCYSGILKTLRCRNEKKHR 240
 Db 181 CQEDSVYVCGPFPRGWNNTIMRNLLGIVLPLIMIVCYSGILKTLRCRNEKKHR 240
 Qy 241 AYRVFTIMIVYFLWTPNIVILLNTQEFFGLSNCESTSOLDOATQVETLGMTMHCCT 300
 Db 241 AYRVFTIMIVYFLWTPNIVILLNTQEFFGLSNCESTSOLDOATQVETLGMTMHCCT 300
 Qy 301 NPIIYAFGEKFRRLSYFFRKHITKRFCKOCPVYRETDGVTSNTNSTGEOEVSAGL 360
 Db 301 NPIIYAFGEKFRRLSYFFRKHITKRFCKOCPVYRETDGVTSNTNSTGEOEVSAGL 360

RESULT 3
 US-09-939-719-7
 ; Sequence 7, Application US/09938719
 ; Patent No. US2002010674A1
 ; GENERAL INFORMATION:
 ; APPLICANT: SAMSON, MICHEL
 ; PARMENTIER, MARC
 ; VASSART, GILBERT
 ; LIBERT, FREDERICK

RESULT 4
 US-09-939-226-7
 ; Sequence 7, Application US/09939226
 ; Patent No. US20020110805A1
 ; GENERAL INFORMATION:
 ; APPLICANT: SAMSON, MICHEL
 ; PARMENTIER, MARC
 ; VASSART, GILBERT

TITLE OF INVENTION: ACTIVE AND INACTIVE CC-CHEMOKINES RECEPTOR AND NUCLEIC ACID MOLECULES ENCODING SAID RECEPTOR
 NUMBER OF SEQUENCES: 17
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Knobbe, Martens, Olson & Bear
 STREET: 620 Newport Center Drive 16th Floor
 CITY: Newport Beach
 STATE: CA
 COUNTRY: U.S.A.
 ZEP: 92660
 COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIP Release #1.0, Version #1.25 (EPO)
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/938,719
 FILING DATE: 24-Aug-2001
 CLASSIFICATION: <Unknown>
 PRIORITY APPLICATION DATA:
 APPLICATION NUMBER: 09/626, 939
 FILING DATE: 27-JULY-2000
 ATTORNEY/AGENT INFORMATION:
 NAME: Altman, Daniel E.
 REGISTRATION NUMBER: 34,115
 REFERENCE/DOCKET NUMBER: <Unknown>
 INFORMATION FOR SEQ ID NO: 7:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 360 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 MOLECULE TYPE: No. 2002010674A1e
 ; SEQUENCE DESCRIPTION: SEQ ID NO: 7:
 US-09-938-719-7

Query Match Score 98.6%; Pred. No. 7.6e-161; Length 360;
 Best Local Similarity 98.3%; Mismatches 2; Indels 4; Gaps 0;
 Matches 354; Conservative 2;

Qy	1	MLSTSRSRFTNTNESEEYTFFDYGACHKFVDKIQGAQLLPPLYSVLFIFGVGN	60
Db	1	MLSTSRSRFTNTNESEEYTFFDYGACHKFVDKIQGAQLLPPLYSVLFIFGVGN	60
Qy	61	MLVVLLINCKKLKCUDTIVLNLAISDLLELTPLWAHSANNEWFGNAMCKLFGLY	120
Db	61	MLVVLLINCKKLKCUDTIVLNLAISDLLELTPLWAHSANNEWFGNAMCKLFGLY	120
Db	61	MLVVLLINCKKLKCUDTIVLNLAISDLLELTPLWAHSANNEWFGNAMCKLFGLY	120
Qy	121	HIGYEGGIFTILLTIDRYLAIVHAWALKARTVTGGVTSVITWLVAFAVGIFTK	180
Qy	121	HIGYEGGIFTILLTIDRYLAIVHAWALKARTVTGGVTSVITWLVAFAVGIFTK	180
Db	121	HIGYEGGIFTILLTIDRYLAIVHAWALKARTVTGGVTSVITWLVAFAVGIFTK	180
Qy	181	CQEDSVYVCGPFPRGWNNTIMRNLLGIVLPLIMIVCYSGILKTLRCRNEKKHR	240
Db	181	CQEDSVYVCGPFPRGWNNTIMRNLLGIVLPLIMIVCYSGILKTLRCRNEKKHR	240
Qy	241	AYRVFTIMIVYFLWTPNIVILLNTQEFFGLSNCESTSOLDOATQVETLGMTMHCCT	300
Db	241	AYRVFTIMIVYFLWTPNIVILLNTQEFFGLSNCESTSOLDOATQVETLGMTMHCCT	300
Qy	301	NPIIYAFGEKFRRLSYFFRKHITKRFCKOCPVYRETDGVTSNTNSTGEOEVSAGL	360
Db	301	NPIIYAFGEKFRRLSYFFRKHITKRFCKOCPVYRETDGVTSNTNSTGEOEVSAGL	360

RESULT 5
 US-09-939-719-7
 ; Sequence 7, Application US/09938719
 ; Patent No. US2002010674A1
 ; GENERAL INFORMATION:
 ; APPLICANT: SAMSON, MICHEL
 ; PARMENTIER, MARC
 ; VASSART, GILBERT
 ; LIBERT, FREDERICK

TITLE OF INVENTION: ACTIVE AND INACTIVE CC-CHEMOKINES RECEPTOR AND NUCLEIC ACID MOLECULES ENCODING SAID RECEPTOR

NUMBER OF SEQUENCES: 17

CORRESPONDENCE ADDRESS:

ADDRESSEE: Knobbe, Martens, Olson & Bear

STREET: 620 Newport Center Drive 16th Floor

CITY: Newport Beach

STATE: CA

COUNTRY: U.S.A.

ZIP: 92660

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PATENTIN Release #1.0, version #1.25 (EPO)

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/939,226

FILING DATE: 24-Aug-2001

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/626,939

FILING DATE: 2000-07-27

ATTORNEY/AGENT INFORMATION:

NAME: Altmann, Daniel E

REGISTRATION NUMBER: 34,115

REFERENCE/DOCKET NUMBER: <Unknown>

INFORMATION FOR SEQ ID NO: 7:

SEQUENCE CHARACTERISTICS:

LENGTH: 360 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: No. US20020110805A1

SEQUENCE DESCRIPTION: SEQ ID NO: 7:

US-09-939-226-7

Query Match 98.6%; Score 1873; DB 10; Length 360;

Best Local Similarity 98.3%; Pred. No. 7.6e-161; Gaps 0;

Matches 354; Conservative 2; Mismatches 2; Indels 4; Gaps 0;

Db 1 MLSTSRSRIRNTNESEGVTTFFDYDGAAPCKKFDVQKIGAQQLLPPPLSVLFIFGFVN 60

Qy 1 MLSTSRSRIRNTNESEGVTTFFDYDGAAPCKKFDVQKIGAQQLLPPPLSVLFIFGFVN 60

Qy 61 MLVVLILINCKKLCKLTDIYLNAISDLFLITLPLWAHSAAAEWGNAMCKLFTGLY 120

Db 61 MLVVLILINCKKLCKLTDIYLNAISDLFLITLPLWAHSAAAEWGNAMCKLFTGLY 120

Qy 121 HIGYEGGIFTILLTIDRYLAIVAHAVEALKARTYTFEGVVTTSVTLVAVFASVPGIIFTK 180

Db 121 HIGYEGGIFTILLTIDRYLAIVAHAVEALKARTYTFEGVVTTSVTLVAVFASVPGIIFTK 180

Qy 181 CQKEDSVYYVCGPYFPRGWNFHFTIMRNLLGVLPLIMIVCYSGILKTLLCRNEKKHR 240

Db 181 CQKEDSVYYVCGPYFPRGWNFHFTIMRNLLGVLPLIMIVCYSGILKTLLCRNEKKHR 240

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

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Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

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Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

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Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

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Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

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Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

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Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

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Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

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Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

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Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

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Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

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Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Db 301 NPITYAFGEKEFRRLVSFEFRKHIXXXFCRKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

Qy 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Db 241 AVRVIIFTIMIVYFLFWTPYNIVLNLNTQEFFGLNSCNESTSQLDAQTVEFTLGTHCC 300

Qy 301 NPITYAFGEKEFRRLVSFEFRKHITKRFCKOCPVFYRETVDGVTSNTPSTGEQEVSAGL 360

P6

Publication No. US20030023044A1
 GENERAL INFORMATION:
 APPLICANT: Ruben, Steven M.
 APPLICANT: Rosen, Craig A.
 TITLE OF INVENTION: HUMAN G-PROTEIN RECEPTOR HGBER32
 NUMBER OF SEQUENCES: 7
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI,
 STREET: 6 Becker Farm Road
 CITY: Roseland
 STATE: New Jersey
 COUNTRY: USA
 ZIP: 07068
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/09/104,792
 FILING DATE:
 CLASIFICATION:
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US/08/461,244
 FILING DATE: 5-JUN-1995
 ATTORNEY/AGENT INFORMATION:
 NAME: Ferraro, Gregory D.
 REGISTRATION NUMBER: 36,134
 REFERENCE/DOCKET NUMBER: 325800-445
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 201-994-1700
 TELEFAX: 201-994-1744
 INFORMATION FOR SEQ ID NO: 3:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 347 amino acids
 TYPE: amino acid
 SPANNINGNESS:
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 MOLECULE ID: 0-9104-792-3

Query Match 96.7%; Score 1838; DB 10; Length 347;
 best Local Similarity 100.0%; Pred. No. 1e-157; Indels 0; Gaps 0;

14 NESGEBEVTPFDYDGAAPCHKFDVQGAQLPPLYSLVFIGFGVNMVLVLLINCKKL
 1 NESGEBEVTPFDYDGAAPCHKFDVQGAQLPPLYSLVFIGFGVNMVLVLLINCKKL 73
 74 KCLTDIYLNLAIISPLFLITPLRAHSANAEVNFGNAMCKLFTGLHYIGYRGIFILL 133
 61 KCLTDIYLNLAIISPLFLITPLWAHSAANEWVNFGNAMCKLFTGLHYIGYRGIFILL 120
 134 LTIDRYLAIVHAVEALKARVTFGVTTSVITWLVAFAVGSKPQEDSVYVCGPY 193
 121 LTIDRYLAIVHAVEALKARVTFGVTTSVITWLVAFAVGSKPQEDSVYVCGPY 180
 194 FPRGWNFHFTIMRNLLGVLPLIMIVCYSGILKTLRCNEKKHRAYRVLFITMIVYF 253
 181 FPRGWNFHFTIMRNLLGVLPLIMIVCYSGILKTLRCNEKKHRAYRVLFITMIVYF 240
 254 LFWTPYNVIVLNNTFQEFLSNCESTSOLDQATOVTETLGMTTHCCINPPIYAFIGEKFKR 313
 241 LFWTPYNVIVLNNTFQEFLSNCESTSOLDQATOVTETLGMTTHCCINPPIYAFIGEKFKR 300
 314 RYLSVFERKHITKRFCKQCPFPYRETVDGYTSTNPSTGQEVSAGL 360
 301 RYLSVFERKHITKRFCKQCPFPYRETVDGYTSTNPSTGQEVSAGL 347

Query Match 82.6%; Score 1568.5; DB 9; Length 344;
 best Local Similarity 95.3%; Pred. No. 1.9e-132; Indels 7; Gaps 5;
 Matches 300; Conservative 3; Mismatches 5; Indels 7;

QY 18 EEVTTFFDYDGAAPCHKFDVQGAQLPPLYSLVFIGFGVNMVLVLLINCKKLCT 71
 Db 1 EEVTTFFDYDGAAPCHKFDVQGAQLPPLYSLVFIGFGVNMVLVLLINCKKLCT 60

QY 78 DIVLNLAISDLPLFLITPLWAHSAANEWVNFGNAMCKLFTGLHYIGYEGGIFTILRID 13
 Db 61 DIVLNLAISDLPLFLITPLWAHSAANEWVNFGNAMCKLFTGLHYIGYEGGIFTILID 12

QY 138 RYLAVHAVFALKARTVTSVVTLYAVASPGIITPKQKEDSVYVCGPYFPRG 19
 Db 121 RYLAVHAVFALKARTVTSVVTLYAVASPGIITPKQKEDSVYVCGPYFPRG 18

QY 198 WNNFHTIMRNLLGVLPLIMIVCYSGILKTLRCNEKKHRAYRVLFITMIVYFLWT 25
 Db 181 WNNFHTIMRNLLGVLPLIMIVCYSGILKTLRCNEKKHRAYRVLFITMIVYFLWT 25

RESULT 8
 US-09-779-879A-9
 Sequence 9, Application US/09779879A
 Patent No. US20048786A1
 GENERAL INFORMATION:
 APPLICANT: Rosen, Craig A.
 APPLICANT: Roschke, Viktor
 APPLICANT: Li, Yi
 APPLICANT: Ruben, Steven, M.
 TITLE OF INVENTION: Human G-protein Receptor (CCR5) HDGNR10
 FILE REFERENCE: 1488.115000A
 CURRENT APPLICATION NUMBER: US/09-779-879A
 CURRENT FILING DATE: 2001-02-09
 PRIOR APPLICATION NUMBER: US 60/181,258
 PRIOR FILING DATE: 2000-02-09
 PRIOR APPLICATION NUMBER: US 60/234,336
 NUMBER OF SEQ ID Nos: 58
 SOFTWARE: PatentIn version 3.0
 SEQ ID NO 9

LENGTH: 344
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-779-879A-9

Query Match 82.6%; Score 1568.5; DB 10; Length 344;
Best Local Similarity 95.3%; Pred. No. 1.9e-133; Mismatches 3; Indels 7; Gaps 3;

Matches 302; Conservative 3; Qry 18 EEVTTFFDYDGA~~PKH~~KDVQI~~G~~AQ~~L~~PPLYSV~~F~~YGNMVLVLINCKKLKC~~L~~T 77
DB 1 EEVTTFFDYDGA~~PKH~~KDVQI~~G~~AQ~~L~~PPLYSV~~F~~YGNMVLVLINCKKLKC~~L~~T 60

Qry 78 D~~I~~YLLNLAISDLFLITLPLWA~~S~~AA~~N~~EVFGNAMC~~K~~L~~T~~G~~H~~Y~~F~~G~~I~~F~~T~~L~~T~~ID 137
DB 61 D~~I~~YLLNLAISDLFLITLPLWA~~S~~AA~~N~~EVFGNAMC~~K~~L~~T~~G~~H~~Y~~F~~G~~I~~F~~T~~L~~T~~ID 120

Qry 138 RYLAVHAFALKARTVTFGVTSVITWLAVFASVPG~~I~~F~~T~~KCQKEDSVYCCPYFPRG 197
DB 121 RYLAVHAFALKARTVTFGVTSVITWLAVFASVPG~~I~~F~~T~~KCQKEDSVYCCPYFPRG 180

Qry 198 WNNFTIMRN~~I~~LGVLPLIMIVCYSGI~~K~~TLLRCRNEKKRHR~~V~~R~~V~~ITMIVYFLWT 257
DB 181 WNNFTIMRN~~I~~LGVLPLIMIVCYSGI~~K~~TLLRCRNEKKRHR~~V~~R~~V~~ITMIVYFLWT 240

Qry 258 PYNIVILNTFQEFFGLSNCESTS~~Q~~DQATQ~~V~~TETLGMTHCC~~T~~NP~~I~~TYAFGEKFRRLS 317
DB 241 PYNIVILNTFQEFFGLSNCESTS~~Q~~DQATQ~~V~~TETLGMTHCC~~T~~NP~~I~~TYAFGEKFR--S 297

Qry 318 VPFRKHTIKRECKQCPV 334
DB 298 LF--HIALG-CRIAPL 310

RESULT 10
US-09-725-285-9
; Sequence 9, Application US/09725285
; Patent No. US20010000241A1

; GENERAL INFORMATION:
; APPLICANT: Li, Yi
; INVENTION: Antibodies to Human G-Protein Receptor HDGNR10
; TITLE OF INVENTION: Human G-Protein Receptor HDGNR10
; TITLE OF INVENTION: (CCR5 Receptor)
; FILE REFERENCE: 148.8.1150003
; CURRENT APPLICATION NUMBER: US/09/725,285
; CURRENT FILING DATE: 2000-11-29
; PRIOR APPLICATION NUMBER: 09/339,912
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: 09/195,662
; PRIOR FILING DATE: 1998-11-18
; PRIOR APPLICATION NUMBER: 08/466,343
; PRIOR FILING DATE: 1995-06-06
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: PatentIn version 3.0
; LENGTH: 329

Qry 198.1150003
; TYPE: PRT
; ORGANISM: Protein
US-09-725-285-9

Query Match 77.5%; Score 1473; DB 10; Length 329;
Best Local Similarity 90.5%; Pred. No. 7e-125; Gaps 4;
Matches 287; Conservative 3; Mismatches 5; Indels 22; Gaps 4;

Qry 18 EEVTTFFDYDGA~~CH~~KDVQI~~G~~AQ~~L~~PPLYSV~~F~~YGNMVLVLINCKKLCLT 77
DB 1 EEVTTFFDYDGA~~CH~~KDVQI~~G~~AQ~~L~~PPLYSV~~F~~YGNMVLVLINCKKLCLT 60

Qry 78 D~~I~~YLLNLAISDLFLITLPLWA~~S~~AA~~N~~EVFGNAMC~~K~~L~~T~~G~~H~~Y~~F~~G~~I~~F~~T~~L~~T~~ID 137
DB 61 D~~I~~YLLNLAISDLFLITLPLWA~~S~~AA~~N~~EVFGNAMC~~K~~L~~T~~G~~H~~Y~~F~~G~~I~~F~~T~~L~~T~~ID 105

Qry 138 RYLAVHAFALKARTVTFGVTSVITWLAVFASVPG~~I~~F~~T~~KCQKEDSVYCCPYFPRG 197
DB 106 RYLAVHAFALKARTVTFGVTSVITWLAVFASVPG~~I~~F~~T~~KCQKEDSVYCCPYFPRG 165

Qry 198 WNNFTIMRN~~I~~LGVLPLIMIVCYSGI~~K~~TLLRCRNEKKRHR~~V~~R~~V~~ITMIVYFLWT 257
DB 166 WNNFTIMRN~~I~~LGVLPLIMIVCYSGI~~K~~TLLRCRNEKKRHR~~V~~R~~V~~ITMIVYFLWT 225

Qry 258 PYNIVILNTFQEFFGLSNCESTS~~Q~~DQATQ~~V~~TETLGMTHCC~~T~~NP~~I~~TYAFGEKFRRLS 317
DB 226 PYNIVILNTFQEFFGLSNCESTS~~Q~~DQATQ~~V~~TETLGMTHCC~~T~~NP~~I~~TYAFGEKFR--S 282

Qry 318 VPFRKHTIKRECKQCPV 334
DB 283 LF--HIALG-CRIAPL 295

RESULT 11

US-09-195-662A-9
Sequence 9, Application US/09195662A
; Patent No. US0020076745A1
; GENERAL INFORMATION:
; APPLICANT: Li, Yi
; APPLICANT: Ruben, Steven, M.
; TITLE OF INVENTION: Human G-Protein Chemokine Receptor HDGNR10 (CCRS Receptor)
; CURRENT APPLICATION NUMBER: 1488 1150002
; FILE REFERENCE: PCT/US1995/06243
; CURRENT FILING DATE: 1998-11-18
; PRIOR APPLICATION NUMBER: 08/466, 343
; PRIOR FILING DATE: 1995-06-06
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO: 9
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Protein
US-09-195-662A-9

Query Match 77.5%; Score 1473; DB 10; Length 329;
Best Local Similarity 90.5%; Pred. No. 7e-125; Indels 22; Gaps 4;
Matches 287; Conservative 3; Mismatches 5;

QY 18 EEVTFDFDYAPCHKFDFKQIGAQOLLPPSLVIFGFGVNMLVVLINCKKLKCLT 77
Db 1 EEVTFDFDYAPCHKFDFKQIGAQOLLPPSLVIFGFGVNMLVVLINCKKLKCLT 60

QY 78 DIYLNLAIISDLFLITLPLWAHSAAANEWFGNAMCKLFTGLYHIGYFGGIFILLITID 137
Db 61 DIYLNLAIISDLFLITLPLWAHSAAANEWFGNAMCKLFTGLYH----- 105

QY 138 RYLAIVHAVFAALKARTVTGFVTVTSLVAVFASPGIITFKCQEDSVYVCGPYFPRG 197
Db 106 RYLAIVHAVFAALKARTVTGFVTVTSLVAVFASPGIITFKCQEDSVYVCGPYFPRG 165

QY 198 WNNFHTIMRNILGLVPLLMIVCGSILKTLRCRNEKKHRRAVIFTIMIVFLEWT 257
Db 166 WNNFHTIMRNILGLVPLLMIVCGSILKTLRCRNEKKHRRAVIFTIMIVFLEWT 225

QY 258 PYNIVILLNTFQEFPGLNSCESTSQDQATOVTETLGMTHCINCINPIYAFGEKFRYRLS 317
Db 226 PYNIVILLNTFQEFPGLNSCESTSQDQATOVTETLGMTHCINCINPIYAFGEKFR--S 282

QY 318 VFFRKHITKRFCKOCPV 334
Db 283 LF--HIALG-CRIAPL 295

RESULT 13
US-09-502-783A-9
; Sequence 9, Application US/09502783A
; Patent No. US200213229A1
; GENERAL INFORMATION:
; APPLICANT: Li, Yi
; APPLICANT: Ruben, Steven M.
; TITLE OF INVENTION: Polynucleotides Encoding Human G-Protein Chemokine Receptor (CCRS Receptor)
; FILE REFERENCE: HDGNR10
; CURRENT APPLICATION NUMBER: 1488 1150006
; CURRENT FILING DATE: 2001-08-23
; PRIOR APPLICATION NUMBER: 08/466, 343
; PRIOR FILING DATE: 1995-06-06
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO: 9
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Protein
US-09-502-783A-9

Query Match 77.5%; Score 1473; DB 10; Length 329;
Best Local Similarity 90.5%; Pred. No. 7e-125; Indels 22; Gaps 4;
Matches 287; Conservative 3; Mismatches 5;

QY 18 EEVTFDFDYAPCHKFDFKQIGAQOLLPPSLVIFGFGVNMLVVLINCKKLKCLT 77
Db 1 EEVTFDFDYAPCHKFDFKQIGAQOLLPPSLVIFGFGVNMLVVLINCKKLKCLT 60

QY 78 DIYLNLAIISDLFLITLPLWAHSAAANEWFGNAMCKLFTGLYHIGYFGGIFILLITID 137
Db 61 DIYLNLAIISDLFLITLPLWAHSAAANEWFGNAMCKLFTGLYH----- 105

QY 138 RYLAIVHAVFAALKARTVTGFVTVTSLVAVFASPGIITFKCQEDSVYVCGPYFPRG 197
Db 106 RYLAIVHAVFAALKARTVTGFVTVTSLVAVFASPGIITFKCQEDSVYVCGPYFPRG 165

QY 198 WNNFHTIMRNILGLVPLLMIVCGSILKTLRCRNEKKHRRAVIFTIMIVFLEWT 257
Db 166 WNNFHTIMRNILGLVPLLMIVCGSILKTLRCRNEKKHRRAVIFTIMIVFLEWT 225

QY 258 PYNIVILLNTFQEFPGLNSCESTSQDQATOVTETLGMTHCINCINPIYAFGEKFRYRLS 317
Db 226 PYNIVILLNTFQEFPGLNSCESTSQDQATOVTETLGMTHCINCINPIYAFGEKFR--S 282

QY 318 VFFRKHITKRFCKOCPV 334
Db 283 LF--HIALG-CRIAPL 295

RESULT 12
US-09-339-912A-9
; Sequence 9, Application US/09339912A
; Patent No. US2002009916A1
; GENERAL INFORMATION:
; APPLICANT: Li, Yi
; APPLICANT: Ruben, Steven, M.
; TITLE OF INVENTION: Antibodies to Human G-Protein Chemokine Receptor HDGNR10
; FILE REFERENCE: 1488-1150003
; CURRENT APPLICATION NUMBER: US/09/339, 912A
; CURRENT FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: 09/195, 662
; PRIOR FILING DATE: 1998-11-18
; PRIOR APPLICATION NUMBER: 08/466, 343
; PRIOR FILING DATE: 1995-06-06
; NUMBER OF SEQ ID NOS: 9
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO: 9
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Protein
US-09-339-912A-9

